Martin Klima

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/6742119/publications.pdf

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18 papers	439 citations	13 h-index	939365 18 g-index
18	18	18	603 citing authors
all docs	docs citations	times ranked	

#	Article	IF	Citations
1	The mycobacterial <i>guaB1</i> gene encodes a guanosine 5′â€monophosphate reductase with a cystathionineâ€Î²â€synthase domain. FEBS Journal, 2022, 289, 5571-5598.	2.2	2
2	Localization of SARS-CoV-2 Capping Enzymes Revealed by an Antibody against the nsp10 Subunit. Viruses, 2021, 13, 1487.	1.5	12
3	Structural basis for hijacking of the host ACBD3 protein by bovine and porcine enteroviruses and kobuviruses. Archives of Virology, 2020, 165, 355-366.	0.9	7
4	Convergent evolution in the mechanisms of ACBD3 recruitment to picornavirus replication sites. PLoS Pathogens, 2019, 15, e1007962.	2.1	26
5	Phosphatidylinositol 4-kinase III \hat{l}^2 (PI4KB) forms highly flexible heterocomplexes that include ACBD3, 14-3-3, and Rab11 proteins. Scientific Reports, 2019, 9, 567.	1.6	17
6	Kobuviral Non-structural 3A Proteins Act as Molecular Harnesses to Hijack the Host ACBD3 Protein. Structure, 2017, 25, 219-230.	1.6	40
7	Metal ionsâ€binding T4 lysozyme as an intramolecular protein purification tag compatible with Xâ€ray crystallography. Protein Science, 2017, 26, 1116-1123.	3.1	7
8	Rational Design of Novel Highly Potent and Selective Phosphatidylinositol 4-Kinase Ill \hat{I}^2 (PI4KB) Inhibitors as Broad-Spectrum Antiviral Agents and Tools for Chemical Biology. Journal of Medicinal Chemistry, 2017, 60, 100-118.	2.9	50
9	Structural analysis of phosphatidylinositol 4-kinase IIIβ (PI4KB) – 14-3-3 protein complex reveals internal flexibility and explains 14-3-3 mediated protection from degradation in vitro. Journal of Structural Biology, 2017, 200, 36-44.	1.3	28
10	Negative charge and membrane-tethered viral 3B cooperate to recruit viral RNA dependent RNA polymerase 3D pol. Scientific Reports, 2017, 7, 17309.	1.6	18
11	Crystal structures of a yeast 14-3-3 protein from <i>Lachancea thermotolerans</i> in the unliganded form and bound to a human lipid kinase PI4KB-derived peptide reveal high evolutionary conservation. Acta Crystallographica Section F, Structural Biology Communications, 2016, 72, 799-803.	0.4	12
12	Structural insights and in vitro reconstitution of membrane targeting and activation of human PI4KB by the ACBD3 protein. Scientific Reports, 2016, 6, 23641.	1.6	81
13	The high-resolution crystal structure of phosphatidylinositol 4-kinase Il \hat{I}^2 and the crystal structure of phosphatidylinositol 4-kinase Il \hat{I}^4 containing a nucleoside analogue provide a structural basis for isoform-specific inhibitor design. Acta Crystallographica Section D: Biological Crystallography, 2015, 71, 1555-1563.	2.5	21
14	Norbornane-based nucleoside and nucleotide analogues locked in North conformation. Bioorganic and Medicinal Chemistry, 2015, 23, 184-191.	1.4	16
15	Inhibition of vacuolar <scp>ATP</scp> ase attenuates the <scp>TRAIL</scp> â€induced activation of caspaseâ€8 and modulates the trafficking of <scp>TRAIL</scp> receptosomes. FEBS Journal, 2013, 280, 3436-3450.	2.2	19
16	T-cell activation triggers death receptor-6 expression in a NF-κB and NF-AT dependent manner. Molecular Immunology, 2011, 48, 1439-1447.	1.0	32
17	Functional analysis of the posttranslational modifications of the death receptor 6. Biochimica Et Biophysica Acta - Molecular Cell Research, 2009, 1793, 1579-1587.	1.9	31
18	Arf and Rho GAP adapter protein ARAP1 participates in the mobilization of TRAIL-R1/DR4 to the plasma membrane. Apoptosis: an International Journal on Programmed Cell Death, 2008, 13, 423-436.	2.2	20